

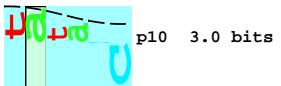
- 1 -

piece 1, NC_000913, yjhU_yjhF+, config: linear, direction: +, begin: 4518318, end: 4518713

... -----] NC_000913.yjhU ... sd



```
{-----> p35-(24)-p10 4518354 Gap 2.4 bits  
-----> p35-p10 4518354 total 5.4 bits
```



The diagram illustrates the structure of a protein domain, likely a nucleic acid binding motif. It features several alpha-helices represented by colored sticks (green, blue, red, orange) and beta-sheets represented by light-colored rectangles. The domain is divided into functional regions: 'sd' (green), 'p35' (blue), '6.1 bits' (red), 'p10' (orange), '1.1 bits' (light blue), 'p10' (cyan), '5.2 bits' (magenta), and another 'p10' region (light blue) with '1.6 bits' (red). The 'p10' regions are associated with short consensus sequences: 'GGGTCCTT' (orange), 'GGGTCCTT' (cyan), 'GGGTCCTT' (magenta), and 'GGGTCCTT' (light blue).

{-----} sd-(8)-ir 4518410 Gap 2.4 bits

p3

ts  p10 2.6 bits

|-----| sd-ir 4518410 yjhU_yjhF+ total 7.2 bits

A small graphic in the bottom right corner featuring a stylized letter 'P'. The letter is composed of several colored segments: a red vertical stroke, a green horizontal bar, a blue diagonal bar, and a red curved segment. It is set against a solid purple rectangular background.

A small image showing a red 'P' and a green vertical bar.

|-----| p35-p10 4518431 total 4.4 bits{-----| p35-(24)-p10 4518442 Gap 2.4 bits-----|

--> p35-p10 45184422 total 6.1 bits
--> p35-(26)-p10 45184444 Gap 3.7 bits

```
|-----| p35-p10 4518447 total 4.3 bits  
|-----| p35-(24)-p10 4518449 Gap 2.4 bits
```

|-----| p35-d10 4518449 total 7.6 bits
|-----| p35-(23)-p10 4518450

|-----| p35-p10 4518456 total 5.0 bits |-----| p35
|-----| 14. |-----|

For more information about the study, please contact Dr. John Smith at (555) 123-4567 or via email at john.smith@researchinstitute.org.

*4518490 *4518500 *4518510 *4518520 *4518530 *4518540 *4518550 *4518560

-fmet - leu - arg - leu - tyr - ser - ile - thr - glu - asn - thr - pro

Arg - Arg - Pro - Ieu - Phe - Pro - His - Cys - Val - Iys - Pro - Gln - Cys - Ser - Gly - Phe - Cys - Val - Ieu - Ieu - Arg - Ile - His - Pro -

[###> orf 25 codons

p10 4.4 bits

p35 2.6 bits

... p10 4.9 bits

